





[illegible][illegible]



RI "MUR" a novel 24000 bp cDNA from human testis  
 RI Cells: 1  
 RI FEBS Lett. 419:215-220(1999).  
 DR EMBL: AF067246; AM28085.1; 1  
 DR PROSITE: P50060; CLAT ADAPTOR M.2.1  
 DR PROSITE: P50094; CLAT ADAPTOR M.2.1  
 SO SEQUENCE 422 AA: 42196 MW: 4825023.1 NC0321

Query Match 33.2% Score 44.5 DB 11 Length 422  
 Best Local Similarity 42.9% Pred. No. 44  
 Matches 9: Conservative 4: Mismatches 2: Indels 6: Gaps 1:

DB 104 RDNFVIVDLELMDPFPQ 124

RESULT 14

ID 061776 PRELIMINARY: PRT: 896 AA.

AC 061776;  
 DT 01-NOV-1998 (TrEMBL) 08, Created  
 DT 01-NOV-1998 (TrEMBL) 08, Last sequence update  
 DT 01-MAY-1999 (TrEMBL) 10, Last annotation update  
 DE HYPOTHETICAL 99.9 KD PROTEIN.  
 OS R28M20.100.  
 OS Arabidopsis thaliana (Mouse-ear cress).  
 CC Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;  
 CC Eudicotyledons; Geraniales; Malvaceae; Malvaceae; Malvaceae;  
 CC Core eudicotyledons; Rosidae; Rosids II; Brassicales; Brassicaceae;  
 CC Arabidopsis.  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RA KIRGER M., MUELLER-ADLER S., ZIPP M., SCHAEFER M., HOEHNEL J.,  
 RA KIRGER M., MUELLER-ADLER S., ZIPP M., SCHAEFER M., HOEHNEL J.,  
 RA KIRGER M., MUELLER-ADLER S., ZIPP M., SCHAEFER M., HOEHNEL J.,  
 PI Submitted (1998-10-15) to EMBL/GenBank/CCDS databases.  
 RN [2]  
 RP SEQUENCE FROM N.A.  
 RA KIRGER M., MUELLER-ADLER S., ZIPP M., SCHAEFER M., HOEHNEL J.,  
 RA KIRGER M., MUELLER-ADLER S., ZIPP M., SCHAEFER M., HOEHNEL J.,  
 RA KIRGER M., MUELLER-ADLER S., ZIPP M., SCHAEFER M., HOEHNEL J.,  
 PI Submitted (1998-10-15) to EMBL/GenBank/CCDS databases.  
 DR EMBL: AL031004; CAA1952.1;  
 DR PIR: P50104; ANP\_receptor; 1.  
 KW Hypothetical protein.  
 SQ SEQUENCE 896 AA: 99925 MW: 985481.090321

Query Match 33.2% Score 44.5 DB 10 Length 898  
 Best Local Similarity 42.9% Pred. No. 44  
 Matches 11: Conservative 4: Mismatches 6: Indels 1: Gaps 1:

DB 680 RDSFVLGLPESCPESRLVPE 701

RESULT 15

ID 035218 PRELIMINARY: PRT: 792 AA.

AC 035218;  
 DT 01-JAN-1998 (TrEMBL) 05, Created  
 DT 01-JAN-1998 (TrEMBL) 05, Last sequence update  
 DT 01-JAN-1998 (TrEMBL) 05, Last annotation update  
 DE CLEAVAGE AND POLYADENYLATION SPECIFICITY FACTOR.  
 GN MOPSF.  
 OS Mus musculus (Mouse).  
 CC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia;  
 CC Eutheria; Rodentia; Sciurotrata; Muridae; Murinae; Mus.  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RA TEVELEV A., SCHATZ D.G.;  
 RA TEVELEV A., SCHATZ D.G.;  
 PI Submitted (1997-03-27) to EMBL/GenBank/CCDS databases.  
 DR EMBL: AF012523; AAB5620.1;  
 DR EMBL: AF012523; AAB5620.1;

52 SEQUENCE 792 AA: 99925 MW: 9770588.090321

Query Match 33.2% Score 44.5 DB 11 Length 792  
 Best Local Similarity 42.9% Pred. No. 44  
 Matches 11: Conservative 4: Mismatches 6: Indels 1: Gaps 1:

Search Completed: January 24, 2000 07:35:41  
 Job time: 221 sec

Thu Jan 27 11:30:43 2000

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